

**Amendments to the Specification:**

Please amend the specification as follows:

Replace paragraph [018] bridging pages 6-7 with the following amended paragraph:

[018] As a rule, the detection sensitivity, [[i.e.,]] a measure of the smallest analyte concentration which can still be detected with confidence, of the test system solid phase R1 and label R2 will be ~~less~~ lower than the detection sensitivity of the test system solid phase R1 and label R3. However, the test system solid phase R1 and label R2 will indicate correct concentration values, even at a relatively high analyte concentration. In other words, the test system "solid phase R1 and label R2" covers the upper measurement range better and the test system "solid phase R1 and label R3" covers the lower measurement range better. By detecting the L1 measurement signal, which is proportional to the formation of the R1-A-R2 complex, separately from the L2-dependent or L1 plus L2-dependent measurement signal, it is possible to broaden the measurement range of the test method according to the invention and/or to recognize a hook effect. For example, if the L1 measurement signal is higher than the highest value on the standard curve and the L2 measurement signal is within the standard curve, also termed calibration curve, that is then a sure sign of a "high-dose hook sample". When a hook effect is detected, the sample is tested once again at an appropriate dilution and the correct analyte concentration is determined.

Replace paragraph [070] on page 27 with the following amended paragraph:

[070] **Fig. 1** shows a diagram of a preferred test method according to the invention ("S" = solid phase). In a first step, solid phase-R1, analyte ("A"; provided it is present in the sample), R2-L1 and R3-L2 (or R3-X) are mixed together and this incubation mixture is incubated until time T1. At time T1, the measurement signal of the label L1 which is contained in the binding complex solid phase-R1-analyte-R2-L1 is

determined. After that, Y-L2 (only in the case of R3-X) is added and the incubation mixture is incubated until time T2. After that, i.e. at time T2, the measurement signal of the label L2 which is present in the binding complex solid phase-R1-analyte-R3-L2 (or - R3-X-Y-L2) is determined. If L1 and L2 are the same labels, preference is given to determining the measurement signal of both the labels L1 and L2 which are present in the binding complexes at time T2. Instead of measuring the measurement signal of the bound labels, it is also possible, in each case or alternatively, to measure the measurement signal of the unbound portion of the label, i.e., the label which is not present in the binding complex, in the incubation mixture.